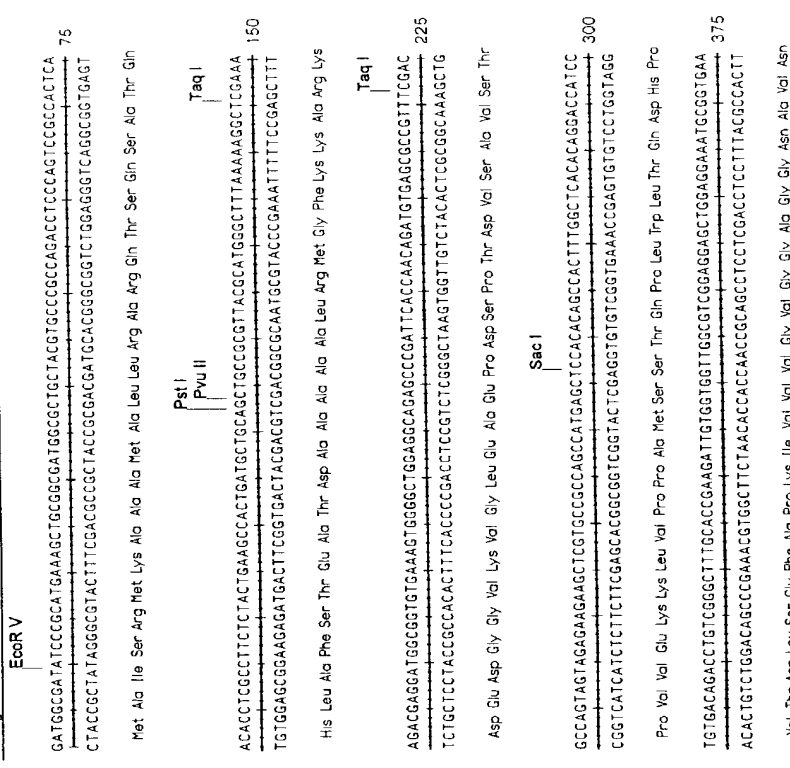


FisZ-m2 consensus2 Map MPD (1 > 1423) Site and Sequence

Enzymes : 50 of 502 enzymes (Filtered)

Settings : Circular, Certain Sites Only, Standard Genetic Code



FisZ-m2 consensus2 Map MPD (1 > 1423) Site and Sequence

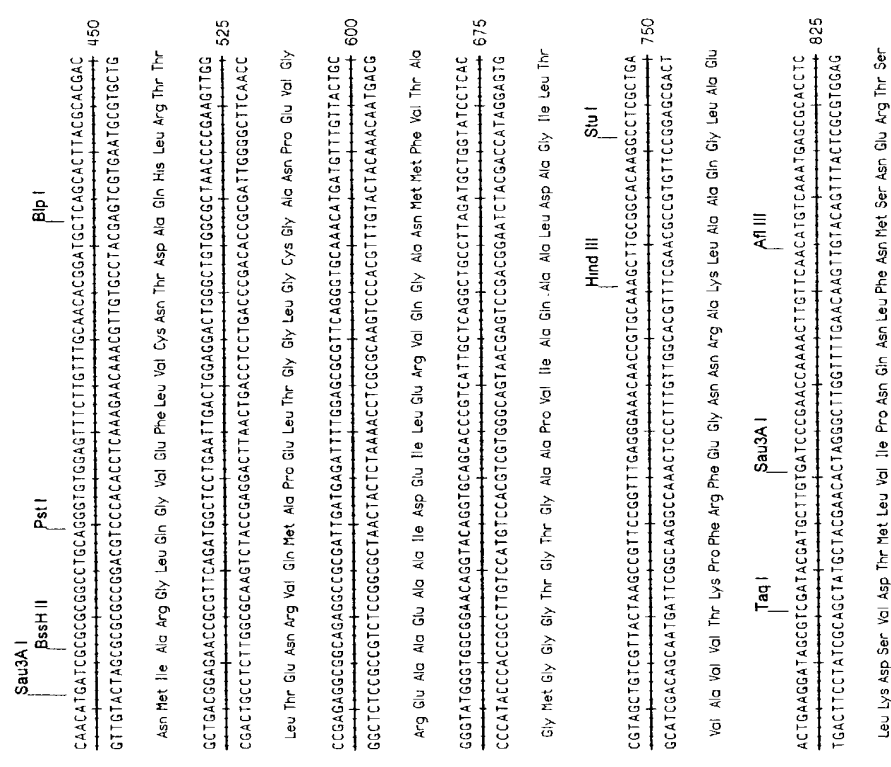


FIG. 1

TTGCCAGTATCAAAAGCGTAAGCAGGGGGAATGACACTAATGACGTGATTGCTTCAGAAATCTCTACAATTGAA
AACCGGTGATAGTTTCGATTTCGTCCCTTACTGTGGATTACTGCACCTAACGAGTGCTTTTAGAGATGTTAAACIT

Cla I Sau3A I
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GTGGCATCGATGTCCTCACGCCCGCGCGTGTGATCGGATTGGTATTATACGGACTGCTTCATCTAGTT
CACCGTAGCTACAGAGGTGGTGGGCGCGACGACTAGCTTAACCATAATATGCTTGACGAAGTATGAATCAA

Cla I
Taq I
Sau3A I

GTGGCATCGATGCTCTCAGCAGACCGCGCGTCTGATCGGATTGGTATTACGGACTGCTTCATCTAGTT
CAGCTAGCTACAGAGTGGCTGGCGGCGACGACTACCTAACCAATAATGCTGACGAAGTATGAATCAA

GTTGTAGCGCATTTCAGAAATGGCGGACAATGCTCTTGGACGGTGTCAAGAACCATTCGGATTGATGGTGT 900
 CAACCTACCTCGGTAAGTCTTACCGCCTGTTACACGAAGACCTGCCACAGTCTCTGTAAAGCCTAAATACCACTA
 Leu Met Asp Ala Phe Arg Met Ala Asp Asn Val Leu Leu Asp Gly Val Lys Asn Ile Ser Asp Leu Met Val Met
 GCTTGGGCTCATTAACTTGACTTTCGGGATGTTCATCGGTATCGGAAATATGGAAAGACGTATGCGGAAG 975
 CGGACCCGAGTAATTTGGAGCTGAAACGCCATCAAGTTTACCGCTAGCTTTTATACCCCTTTGGCTACTACCCCTTC

ISBN

TGGAGAGCCGGATGGAGAGAA TGGGCTCTGGTGCTGCTGAAGATGCCATGGCGAACCCCTCTTC TGGGGTGATAT 1050

[illegible]

Sau3A I

TTCGATTAGGAGGCCAAGGCATGATCGTTAATATCAGGGAGGCTCGACCTACGCGTATTGAGTTCATGA 1125

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Saul3A1

TaqI TaqI

GGCTGCTGACGGTGTGACCGGGAACTTCATGATCCACACGCCAACATCATCTTCGGTTCGACTTCGACGACTC
1200

Protein

Affiliations:
MILU I

GC TGG GCG GCA AGC TAC CGG CTC CGT GGT TGC CAC TGG TAT TGC CGA CCC GCA CAG TTA TAG AAC CCG GAT G 1275

Leu, Cys, Gly, Leu, Arg, Val, Ser, Val, Val, Ala, Thr, Gly, Ile, Ala, Asp, Pro, Asp, Lys, Leu •

	Source Organism (organelle)	GenBank Accession No.
SEQ ID NO: 11	<i>Agrobacterium tumefaciens</i>	O30992
SEQ ID NO: 12	<i>Sinorhizobium meliloti</i>	P30327
SEQ ID NO: 13	<i>Bartonella clarridgeiae</i>	AAD31718
SEQ ID NO: 14	<i>Rickettsia prowazekii</i>	Q9ZCQ3
SEQ ID NO: 15	<i>Caulobacter crescentus</i>	P52976
SEQ ID NO: 16	<i>Cyanidioschyzon merolae</i> (mt)	BAA85115
SEQ ID NO: 4	<i>Phytophthora infestans</i> -mt2	this invention
SEQ ID NO: 17	<i>Mallomonas splendens</i> (mt)	AAF35432
SEQ ID NO: 2	<i>Phytophthora infestans</i> -mt1	this invention
SEQ ID NO: 18	<i>Gentiana lutea</i> (cp)	T51088
SEQ ID NO: 19	<i>Nicotiana tabacum</i> (cp, 2-1)	T51087
SEQ ID NO: 20	<i>Arabidopsis thaliana</i> (cp, 2-1)	T49028
SEQ ID NO: 21	<i>Physcomitrella patens</i> (cp, 1)	T51089
SEQ ID NO: 22	<i>Physcomitrella patens</i> (cp, 2)	T51090
SEQ ID NO: 23	<i>Guillardia theta</i> (cp)	CAB40398
SEQ ID NO: 24	<i>Mallomonas splendens</i> (cp)	AAF35433
SEQ ID NO: 25	<i>Anabaena PCC7120</i>	CAA83241
SEQ ID NO: 26	<i>Synechocystis PCC6803</i>	P73456
SEQ ID NO: 27	<i>Arabidopsis thaliana</i> (cp, 1-1)	Q42545
SEQ ID NO: 28	<i>Pisum sativum</i> (cp)	T06774
SEQ ID NO: 29	<i>Nicotiana tabacum</i> (cp, 1-3)	CAB89287
SEQ ID NO: 30	<i>Nicotiana tabacum</i> (cp, 1)	CAB41987
SEQ ID NO: 31	<i>Nicotiana tabacum</i> (cp, 1-1)	CAB89286
SEQ ID NO: 32	<i>Nicotiana tabacum</i> (cp, 2)	AAF23770

<u>Bacterial FtsZ</u>	1	50
SEQ ID NO: 11	PRITVFGVGGGGGNAVNMMITVGLQGVDFVVANTDAQALMT..KADRVQLGVNVTEGL	
SEQ ID NO: 12	PRITVFGVGGGGGNAVNMMITAGLQGVDFVVANTDAQALMT..KAERIIQMGVAVTEGL	
SEQ ID NO: 13	PRITVFGVGGGGGNAVNMMINAGLQGVDFVVANTDAQALAMS..KAERVQLGAAVTEGL	
SEQ ID NO: 14	PTITVFGVGGAGSNAVNMMIHANLQGANFVVANTDAQSLEHS..LCINKIQLGVSTTRGL	
SEQ ID NO: 15	PRIVVFGVGGAGGNAVNMMIEAGLEGVEFVVANTDAQQLQFA..KTDRIQLGVQITQGL	
<u>Mitochondrial FtsZ</u>		
SEQ ID NO: 16	PRIMVVGVGAGGNAVNMMIASSLPGEFLVANTDAQALKMS..LCPNRIQLGASLTEGL	
SEQ ID NO: 4	PKIVVVGVGAGGNAVNMMIARGLQGVFLVCNTDAQHLRTT..LTENRVQMAPELTGGL	
SEQ ID NO: 17	PKICVFGVGGGGCNAVNMMIARKLSGVEFVCANTDAQHLSTC..LTENKLQLGKESTQGL	
SEQ ID NO: 2	AS.....QLEGVEFIVANTDCQALGRS..LAPHKITLGKDITKGL	
<u>Chloroplast FtsZ</u>		
SEQ ID NO: 18	AKIKVVGVGGGGSNAVNRMIESAMKGVEFWIVNTDVQAIKMSPVYLENRLQIGQELTRGL	
SEQ ID NO: 19	AKIKVVGVGGGGSNAVNRMIESSMKGVEFWIVNTDIQAMRMSPVAAEQRLPIGQELTRGL	
SEQ ID NO: 20	ARIKVIQVGGGGSNAVNRMIESEMSGVEFWIVNTDIQAMRMSPVLPDNRLQIGKELTRGL	
SEQ ID NO: 21	AKIKVIQVGGGGSNAVNRMLESEMQGVEFWIVNTDAQAMALSPVPAQNRLQIGQKLTRGL	
SEQ ID NO: 22	AKIKVIQVGGGGSNAVNRMLESEMQGVEFWIVNTDAQAMALSPVPAQNRLQIGQKLTRGL	
SEQ ID NO: 23	CVIKVIQVGGGGSNAVNRMVG.GVEGVEFWSINTDAQALSRS..LAPNTCNIGAKLTRGL	
SEQ ID NO: 24GVELWVVNTDAQALSRS..SAKRRLNIGKVLRSGL	
SEQ ID NO: 25	ANIKVIQVGGGGSNAVNRMIESDVSGVEFWSINTDAQALTLA..GAPSRLQIGQKLTRGL	
SEQ ID NO: 26	AKIKVIQVGGGGSNAVNRMIASGVTGIDFWAINTDSQALTNT..NAPDCIQIGQKLTRGL	
SEQ ID NO: 27	ARIKVIQVGGGGSNAVNRMISGLQSVDFYAINNTDSQALLQFSA..ENPLQIGELLTRGL	
SEQ ID NO: 28	AKIKVVGIGGGGSNAVNRMIGSGLQGVDFYAINNTAQALLHSAA..ENPIKIGELLTRGL	
SEQ ID NO: 29	AKIKVIQVGGGGSNAVNRMIGSGLQGVDFYAINNTAQALLQSAA..ENPLQIGELLTRGL	
SEQ ID NO: 30	AKIKVIQVGGGGSNAVNRMIGSGLQGVDFYAINNTAQALLQSAA..ENPLQIGELLTRGL	
SEQ ID NO: 31	AKIKVVGVGGGGSNAVNRMIGSGLQGVDFYAVNTAQALLQSTV..ENPIQIGELLTRGL	
SEQ ID NO: 32	AKIKVVGVGGGGSNAVNRMIGSGLQGVDFYAVNTAQALLQSTV..ENPIQIGELLTRGL	

FIG. 2

<u>Bacterial FtsZ</u>	60	110
SEQ ID NO: 11	GAGSQPEVGRAAAEECIDDEIIDHLNGTHMCFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 12	GAGSQPEVGRAAAEECIDDEIIDHLQGTHMCFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 13	GAGALPEVGRAAADECIDEIIDHLADSHMVFITAGMGGGTGTGAAPVVAANAAREKGILTV	
SEQ ID NO: 14	GAGASPEVGALAAQESENEIRSSLENSNMVFITAGMGGGTGTGSAPIIARIAKELGILTV	
SEQ ID NO: 15	GAGAHPEVGMSAAEESFPEIGEHLGAMHVFITAGMGGGTGTGAAPIIAKCARERGILTV	
<u>Mitochondrial FtsZ</u>		
SEQ ID NO: 16	GAGARPDIGRAAAEEAYETLKREFRGVHLLFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 4	GCGANPEVGREAAEAIDEILERVQGANMMFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 17	GCGANPESGRRAAEESKEEIIARYIADANMVFITAGMGGGTGTGAAPVVAEVCMEKDILTV	
SEQ ID NO: 2	GAGSKPELGKRSAEQQKVDIQRMLQDSNMLFITGGMGGGTCTGAAPVVASVARELGILTV	
<u>Chloroplast FtsZ</u>		
SEQ ID NO: 18	GAGGNPDIGMNAAKESKEAIEEAVYGADMVFTAGMGGGTGTGGAPVIAGIAKSMGILTV	
SEQ ID NO: 19	GAGGNPDIGMNAANESKQAEIEEAVYGADMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 20	GAGGNPEIGMNAARESKEVIEEALYGS DMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 21	GAGGNPEIGCSAAEESKAMVEEALRGADMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 22	GAGGNPEIGCSAAEESKAMVEEALRGADMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 23	GAGGNPEIGRKAEEESRDLIAEAVSAGDLVFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 24	GAGGNPAIGAKAAEESREEIMAVVKNADLVFVITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 25	GAGGNPAIGQKAAEESRDEIATALEGADLVFITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 26	GAGGNPAIGQKAAEESRDEIARSLEGLDLVFITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 27	GTGGNPLLGEQAAEESKDAIANALKGS DLVFITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 28	GTGGNPLLGEQAAEESKEAIIANALKGS DLVFITAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 29	GTGGNPLLGEQAAEESKEAIIANSLKGS DMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 30	GTGGNPLLGEQAAEESKEAIIANSLKGS DMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 31	GTGGNPLLGEQAAEESKEHIIANALKGS DMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
SEQ ID NO: 32	GTGGNPLLGEQAAEESKEHIIANALKGS DMVFTAGMGGGTGTGAAPVVAQAARNKGILTV	
<u>Bacterial FtsZ</u>	120	170
SEQ ID NO: 11	GVVTKPFHFEGGRRMRLAEQEGIEELQKSVDTLIVIPNQNLFRANDKTTFADAFAMADQV	
SEQ ID NO: 12	GVVTKPFHFEGGRRMRLADQGISDLQKSVDTLIVIPNQNLFRANDKTTFADAFAMADQV	
SEQ ID NO: 13	GVVTKPFQFEGARRMKTAEAGIEELQKSVDTLIVIPNQNLFRANDKTTFADAFAMADQV	
SEQ ID NO: 14	GVVTKPFHFEGGHRMKTADKGLIELQQFVDTLIVIPNQNLFRANDKTTFADAFAMADQV	
SEQ ID NO: 15	GVVTKPFHFEGRRMRLADSGIQELQRYVDTLIVIPNQNLFRANDKTTFADAFAMADQV	
<u>Mitochondrial FtsZ</u>		
SEQ ID NO: 16	AVVTKPFHFEGMIRMKTAEQGIVELTEHVD TMLVIPNQNLFKVASPRTSFLDAFRLADHV	
SEQ ID NO: 4	AVVTKPFRFEGNNRAKLAAQGLAELKDSVD TMLVIPNQNLFNMSNERTSLMDAFRMADNV	
SEQ ID NO: 17	AVVTKPFSFEGKRRRLANEGIRSLERVD TLIIPNQNLFKLINASTSMADAFGLADDI	
SEQ ID NO: 2	GVVSTPFRSEGNRTRLANAGVKELAKYVD TLIIVPNQNLALADKSTTMLEAFRYADDV	
<u>Chloroplast FtsZ</u>		
SEQ ID NO: 18	GIVTTPFSFEGRRRAVQAQEGIAALRDND TMLVIPNDKLLTAVSPSTPVTEAFNLADDI	
SEQ ID NO: 19	GIVTTPFSFEGRRRAVQAQEGIAALREND TMLVIPNDKLLTAVSPSTPVTEAFNLADDI	
SEQ ID NO: 20	GIATTPFSFEGRRRTVQAQEGLASLRDND TMLVIPNDKLLTAVSQSTPVTEAFNLADDI	
SEQ ID NO: 21	GIVTTPFAFEGRRRAVQAHEGIAALKNNVD TMLVIPNNKLLTAVAQSTPVTEAFNLADDI	
SEQ ID NO: 22	GIVTTPFAFEGRRRSVQAHEGIAALKNNVD TMLVIPNNKLLTAVAQSTPVTEAFNLADDI	
SEQ ID NO: 23	GVVTKPFAFEGKRRMQQANDAILNLRNKVD TMLVIVVSNKLLQIVPDNTPLQDAFVSADDI	
SEQ ID NO: 24	GVVTKPFGFEGKRRMQQARNAILKMDKVD TMLVIVVSNKLLKIVPDNTPLTEAFVADDI	
SEQ ID NO: 25	GVVTRPFFVFEGRRRRSQAQEGIEGLKSRVD TMLVIPNNKLLLEVIPEQTPVQEAFLYADDV	
SEQ ID NO: 26	GIVTRPFTFEGRRRAKQAEEGINALQSRVD TMLVIPNNQLLSVIPAEQTPVQEAFLYADDI	
SEQ ID NO: 27	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	
SEQ ID NO: 28	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	
SEQ ID NO: 29	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	
SEQ ID NO: 30	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	
SEQ ID NO: 31	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	
SEQ ID NO: 32	GVVTPPFSFEGRRKSLQALEAIEKLQKNVD TMLVIPNDRLLDIADEQTPVQEAFLYADDV	

FIG. 2

<u>Bacterial FtsZ</u>		180	230
SEQ ID NO: 11	LYSGVACITDLMVKEGLINLDFADVRSVMREMARPMGMTGE....	ASGPARAMQAAEAAI	
SEQ ID NO: 12	LYSGVACITDLMVKEGLINLDFADVRSVMREMGGRAMMTGE....	ASGEGRAMAAEAAI	
SEQ ID NO: 13	LYSGVASITDLMIKEGLINLDFADVRSVMHEMGGRAMMTGE....	ASGDGRALAAEAAI	
SEQ ID NO: 14	LHAGVRGVTDLMIMPGLINLDFADIKAVMSEMKGAMMTGE....	DSGEDRAIKAAESAI	
SEQ ID NO: 15	LHSGVRSITDLMVLPGLINLDFADVRTVMTEMGKAMMTGE....	GTAEDRALMAAQNAI	
<u>Mitochondrial FtsZ</u>			
SEQ ID NO: 16	LYSGVRSITDLMTPVGLINLDFADVRSVVREMGGRAMMGSGEVEMEAGNEERAIRASEAAI		
SEQ ID NO: 4	LLDGVKNISDLMVMPLINLDFADVQSVQMGNAMMGSGEAD....	GENRALRAAEDAL	
SEQ ID NO: 17	LLAGVKSITDLMVRPGLINLDFADVRTVMSGMGHAIMGTGQAE....	GEDRAIRAANDAL	
SEQ ID NO: 2	LLEGVKGVTDLIVRPGLINL.....		
<u>Chloroplast FtsZ</u>			
SEQ ID NO: 18	LRQGVRGISDIITIPGLVNVDFAVRAIMANAGSSLMGIGT....	ATGKTRARDAALNAI	
SEQ ID NO: 19	LRQGVRGISDIITIPGLVNVDFAVRAIMANAGSSLMGIGT....	ATGKTRARDAALNAI	
SEQ ID NO: 20	LRQGVRGISDIITIPGLVNVDFAVRAIMANAGSSLMGIGT....	ATGKSRARDAALNAI	
SEQ ID NO: 21	LRQGVRGISDIITVPGLVNVDFAVRAIMANAGSSLMGIGT....	ATGKSRAREAAISAI	
SEQ ID NO: 22	LRQGVRGISDIITVPGLVNVDFAVRAIMANAGSSLMGIGT....	ATGKSKAREAAISAI	
SEQ ID NO: 23	LRQGVVGISEIIVRPGLINVDFAVRSVMADAGSALMGIGT....	GSGKTRAQDAAVAAI	
SEQ ID NO: 24	LRQGVVGITEIIVKPLVNVDFAVRAVMADAGSALMGIGH....	GKGKNAKDAALSAI	
SEQ ID NO: 25	LRQGVQGISDIITIPGLVNVDFAVRAVMADAGSALMGIGV....	SSGKSRAREAAIAAI	
SEQ ID NO: 26	LRQGVQGISDIITIPGLVNVDFAVRAVMADAGSALMGIGV....	GSGKSRAREAAATAAI	
SEQ ID NO: 27	LRQGVQGISDIITIPGLVNVDFAVKAVMKDSGTAMLGVG....	SSSKNRAEEAAEQAT	
SEQ ID NO: 28	LRQGVQGISDIITIPGLVNVDFAVKAVMKDSGTAMLGVG....	SSSKNRAEEAAEQAT	
SEQ ID NO: 29	LRQGVQGISDIITIPGLVNVDFAVKAVMKDSGTAMLGVG....	SSSKNRAEEAAEQAT	
SEQ ID NO: 30	LRQGVQGISDIITIPGLVNVDFAVKAVMKDSGTAMLGVG....	SSSKNRAEEAAEQAT	
SEQ ID NO: 31	LCQGVQGISDIITIPGLVNVDFAVKAIMKDSGTAMLGVG....	SSSRNRAEEAAEQAT	
SEQ ID NO: 32	LCQGVQGISDIITIPGLVNVDFAVKAIMKDSGTAMLGVG....	SSSRNRAEEAAEQAT	
<u>Bacterial FtsZ</u>		240	290
SEQ ID NO: 11	ANPLLD.ETSMKGAQGLLISITGGRDLTLFEVDEAAATRIREEVDP.	DANII LGATFDEAL	
SEQ ID NO: 12	ANPLLD.ETSMKGAQGLLISITGGRDLTLFEVDEAAATRIREEVDP.	DANII LGATFDEEL	
SEQ ID NO: 13	ANPLLD.DTSMRGARGLLISITGGRDMLTLFEVDEAAANRIREEVDA.	DANVIFGAIIDDES	
SEQ ID NO: 14	SNPLLD.HSSMCGARGVLINITGGPDMTLFEVDNAANRIREEVDNIDANI	IFGSTFNPEL	
SEQ ID NO: 15	ANPLLD.EVSLKGAKAVLVNVTGGM DMTLFEVDEAANAISDQVDP.	EANII FGAAFDPSL	
<u>Mitochondrial FtsZ</u>			
SEQ ID NO: 16	CNPLLD.ETSLRGARGVLVNITGGTDMTLFEIDAAANRIREQVDP.	DANII FGSAFDASM	
SEQ ID NO: 4	ANPLLG.DISIKDAKMIVNITGGSDTLTLFEVDEAAERVTRERLDDPHANII	IFGSTFDDSL	
SEQ ID NO: 17	NNPLLGDFSVRSAKGMVLNITGGKDLTLFEVDAAAQRITSEIEDEDANVIFGSS	FDES	
SEQ ID NO: 2		
<u>Chloroplast FtsZ</u>			
SEQ ID NO: 18	QSPLLD..IGIERATGIVWNITGGSDTLTLFEVNAAA EVIYDLVDP.	SANLIFGAVVDPSL	
SEQ ID NO: 19	QSPLLD..IGIERATGIVWNITGGSDTLTLFEVNAAA EVIYDLVDP.	SANLIFGAVIDPSI	
SEQ ID NO: 20	QSPLLD..IGIERATGIVWNITGGSDTLTLFEVNAAA EVIYDLVDP.	TANLIFGAVVDPAL	
SEQ ID NO: 21	QSPLLD..VGIERATGIVWNITGGSDMTLFEVNAAA EVIYDLVDP.	NANLIFGAVVDEAL	
SEQ ID NO: 22	QSPLLD..VGIERATGIVWNITGGSDMTLFEVNAAA EVIYDLVDP.	NANLIFGAVVDEAL	
SEQ ID NO: 23	SSPLLD..FPIEKARGIVFNITGGQDMTLHEINSAA EVIYEAVDS.	NANII FGALVDDNM	
SEQ ID NO: 24	SSPLLD..FPITRAKGIVFNIVGSDMSLQEI NAAA EVIYENVDP.	DANII FGAMVDDKM	
SEQ ID NO: 25	SSPLLE..CSIEGARGVFNITGGSDTLTLHEVNAAA EIIYEVVDP.	NANII FGAVIDDR	
SEQ ID NO: 26	SSPLLE..SSIQGAQGVFNVTGGTDLTLHEVNVA AEI IYEVVDA.	DANII FGAVIDDR	
SEQ ID NO: 27	LAPLIG..SSIQSATGVVYNITGGKDITLQEVNRVSQVVTSLADP.	SANII FGAVVDDRY	
SEQ ID NO: 28	LAPLIG..SSIQSATGVVYNITGGKDITLQEVNRVSQVVTSLADP.	SANII FGAVVDDRY	
SEQ ID NO: 29	LAPLIG..SSIQSATGVVYNITGGKDITLQEVNRVSQVVTSLADP.	SANII FGAVVDERY	
SEQ ID NO: 30	LAPLIG..SSIQSATGVVYNITGGKDITLQEVNRVSQVVTSLADP.	SANII FGAVVDERY	
SEQ ID NO: 31	LAPLIG..LSIQSATGVVYNITGGKDITLQEVNKVSQVVTSLADP.	SANII FGAVVDERY	
SEQ ID NO: 32	LAPLIG..SSIQSATGDVYNITGGKDITLQEVNKVSQVVTSLADP.	SANII FGAVVDERY	

FIG. 2

<u>Bacterial FtsZ</u>	300
SEQ ID NO: 11	E.GLIRVSVVATGI
SEQ ID NO: 12	E.GLIRVSVVATGI
SEQ ID NO: 13	E.GVIRVSVVATGI
SEQ ID NO: 14	K.GIIRVSVVATGI
SEQ ID NO: 15	E.GVIRVSVVATGM
<u>Mitochondrial FtsZ</u>	
SEQ ID NO: 16	Q.GRLRVSVLATGI
SEQ ID NO: 4	G.GKLRVSVVATGI
SEQ ID NO: 17	Q.GSIRVSIVATGI
SEQ ID NO: 2
<u>Chloroplast FtsZ</u>	
SEQ ID NO: 18	C.GQVSITLIATGF
SEQ ID NO: 19	S.GQVSITLIATGF
SEQ ID NO: 20	S.GQVSITLIATGF
SEQ ID NO: 21	H.GQVSITLIATGF
SEQ ID NO: 22	H.DQISITLIATGF
SEQ ID NO: 23	EN.EISITVVATGF
SEQ ID NO: 24	TSGEVSITVLATGF
SEQ ID NO: 25	Q.GEVKITVIATGF
SEQ ID NO: 26	Q.GEMKITVIATGF
SEQ ID NO: 27	.TGEIHVTIIATGF
SEQ ID NO: 28	.TGEIHVTIIATGF
SEQ ID NO: 29	.NGEIHVTIIATGF
SEQ ID NO: 30	.NGEIHVTIIATGF
SEQ ID NO: 31	.NGEIQVTIIATGF
SEQ ID NO: 32	.NGEIQVTIIATGF

FIG. 2

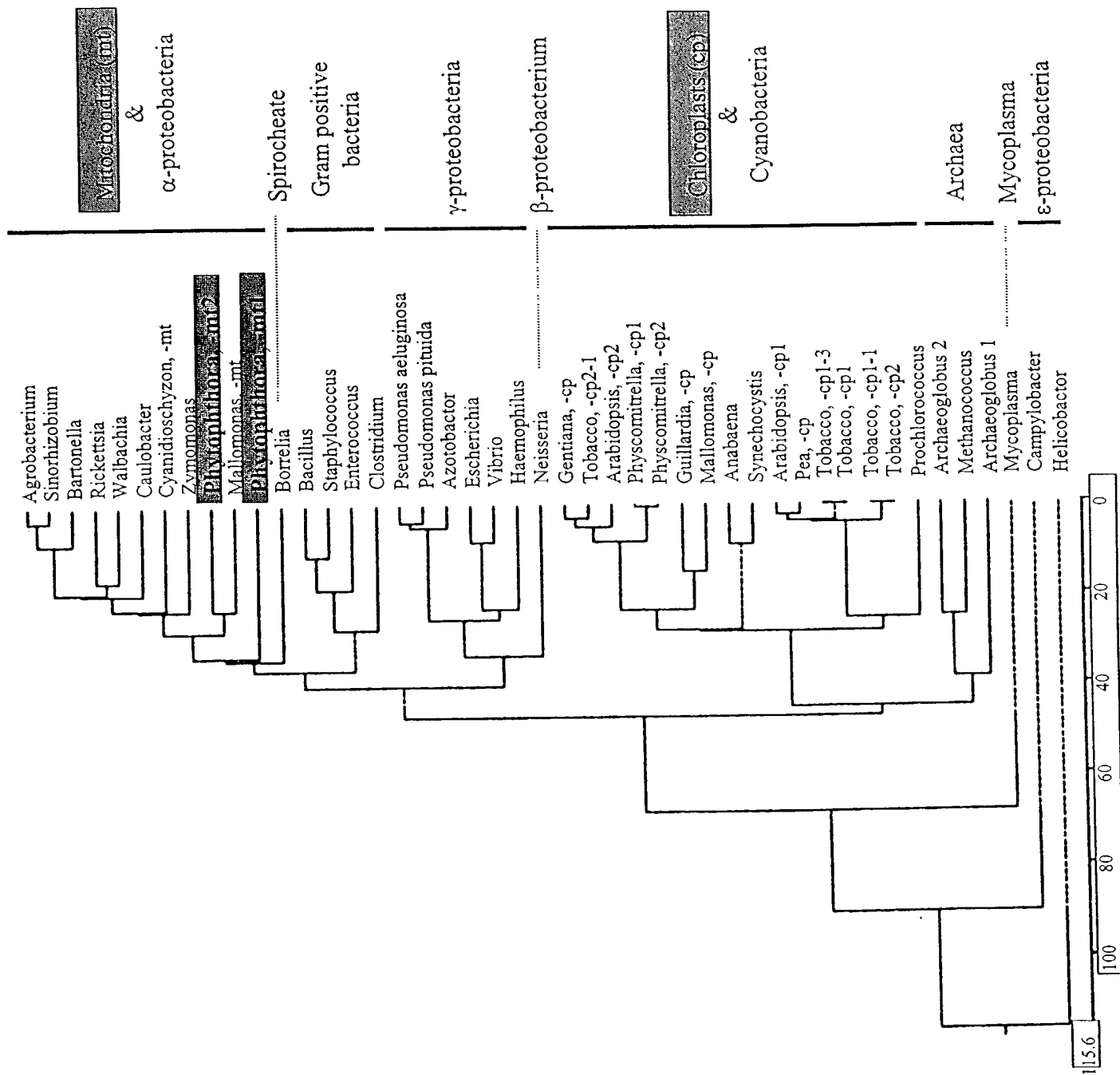


FIG. 3

115.6 100 80 60 40 20 0